

Shin Lin Chen

/633

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/484,064

484, 964

DATE: 05/14/2001
 TIME: 11:03:55

Input Set : A:\Utsh248.txt
 Output Set: N:\CRF3\05142001\I484064.raw

4 <110> APPLICANT: YEH, EDWARD T.H.
 W--> 5 <120> TITLE OF INVENTION: COMPOSITIONS AND USES FOR A NOVEL CELL-DEATH-PROTECTING
 W--> 6 PROTEIN
 9 <130> FILE REFERENCE: UTSH:248
 W--> 10 <140> CURRENT APPLICATION NUMBER: 09/484,964
 C--> 12 <141> CURRENT FILING DATE: 2000-01-18
 12 <150> PRIOR APPLICATION NUMBER: 08/964,162
 13 <151> PRIOR FILING DATE: 1997-11-04
 15 <150> PRIOR APPLICATION NUMBER: 60/030,302
 16 <151> PRIOR FILING DATE: 1996-11-05
 18 <160> NUMBER OF SEQ ID NOS: 16
 19 <170> SOFTWARE: PatentIn Ver. 2.0

Does Not Comply
 Corrected Diskette Needed

ERRORED SEQUENCES

249 <210> SEQ ID NO: 11
 250 <211> LENGTH: 9
 251 <212> TYPE: PRT
 252 <213> ORGANISM: Influenza virus
 254 <400> SEQUENCE: 11
 255 Tyr Pro Tyr Asp Val Pro Asp Tyr Ala
 256 1 5
 258 <210> 12
 259 <211> LENGTH: 4
 260 <212> TYPE: PRT
 261 <213> ORGANISM: Homo sapiens
 W--> 262 <210> SEQ ID NO:
 E--> 263 <400> SEQUENCE: 12
 264 His Ser Thr Val
 265 1
 E--> 267 <210> SEQ ID NO: 13
 324 <210> SEQ ID NO: 16
 325 <211> LENGTH: 30
 326 <212> TYPE: PRT
 327 <213> ORGANISM: Human
 329 <400> SEQUENCE: 16
 330 Val Gln Asp Leu Ala Gln Leu Val Glu Glu Ala Thr Gly Val Pro Leu
 331 1 5 10 15
 332 Pro Phe Gln Lys Leu Ile Phe Lys Gly Lys Ser Leu Lys Glu
 333 20 25 30
 E--> 340 Utsh238.txt
 E--> 341 -1-

Delete extra bracket (<)

Delete extraneous material from the end
 of the file. It is causing processing errors.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/484,064

DATE: 05/14/2001
TIME: 11:03:56

Input Set : A:\Utsh248.txt
Output Set: N:\CRF3\05142001\I484064.raw

L:5 M:283 W: Missing Blank Line separator, <120> field identifier
L:10 M:283 W: Missing Blank Line separator, <140> field identifier
L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:26 M:283 W: Missing Blank Line separator, <220> field identifier
L:29 M:283 W: Missing Blank Line separator, <400> field identifier
L:96 M:283 W: Missing Blank Line separator, <220> field identifier
L:100 M:283 W: Missing Blank Line separator, <220> field identifier
L:104 M:283 W: Missing Blank Line separator, <220> field identifier
L:108 M:283 W: Missing Blank Line separator, <220> field identifier
L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:151 M:283 W: Missing Blank Line separator, <220> field identifier
L:157 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:212 M:283 W: Missing Blank Line separator, <220> field identifier
L:241 M:283 W: Missing Blank Line separator, <220> field identifier
L:263 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:263 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:11 differs:12
L:267 M:214 E: (33) Seq.# missing, SEQ ID NO:12
L:340 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:340 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:341 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:16
L:341 M:252 E: No. of Seq. differs, <211>LENGTH:Input:30 Found:31 SEQ:16